

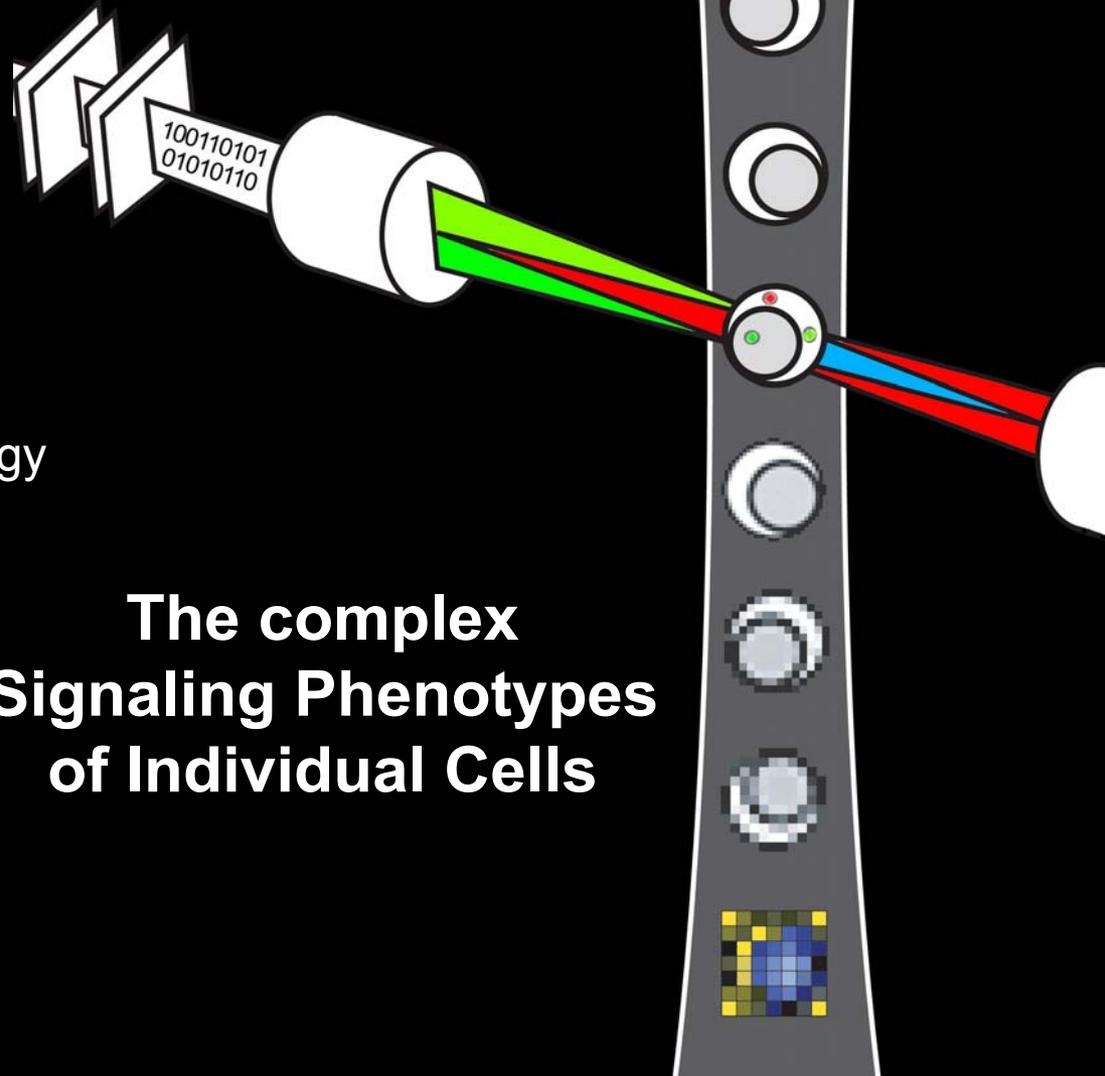
Single Cell Proteomics: A Challenge of Knowing Too Much.

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The complex
Signaling Phenotypes
of Individual Cells



Early Single Cell Proteomics Innovators



Len Herzenberg - - Argon laser flow sorter 1972
- placed an argon laser onto their sorter and successfully did high speed sorting - Coined the term **F**luorescence **A**ctivated **C**ell **S**orting (FACS)



Mack Fulwyler - Coulter Electronics manufactured the TPS-1 (Two parameter sorter) in 1975 which could measure forward scatter and fluorescence using a 35mW argon laser.

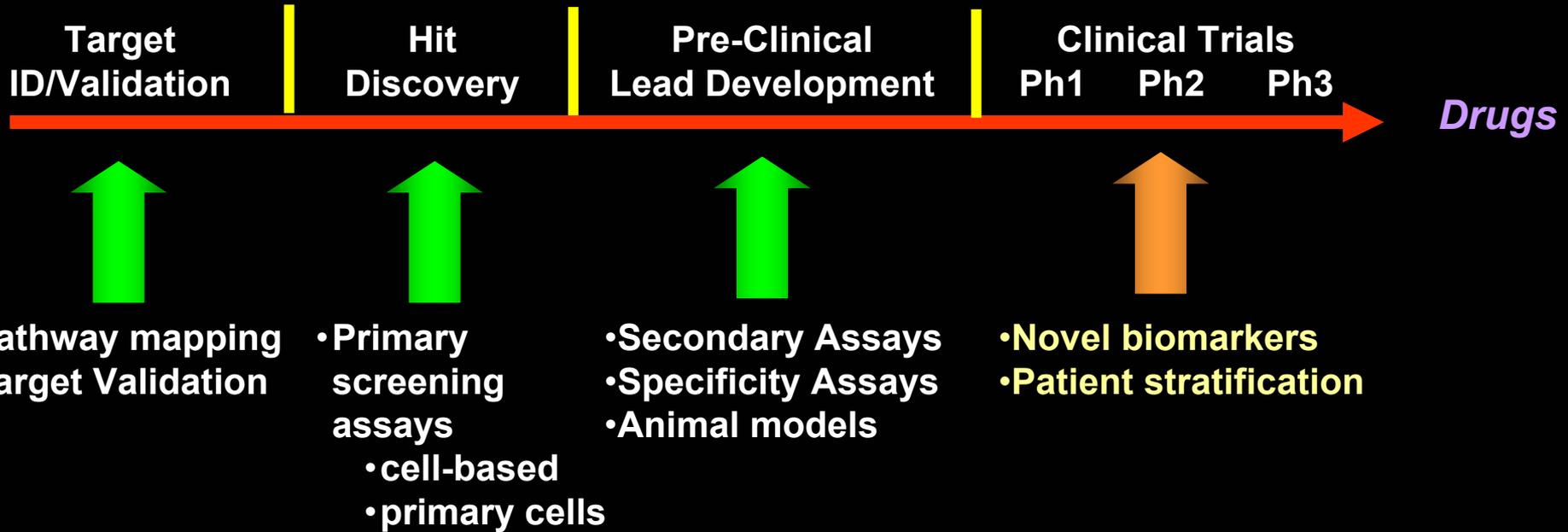


Howard Shapiro: Block instruments (1973-76) - a series of multibeam flow cytometers that did differentials and multiple fluorescence excitation and emission.

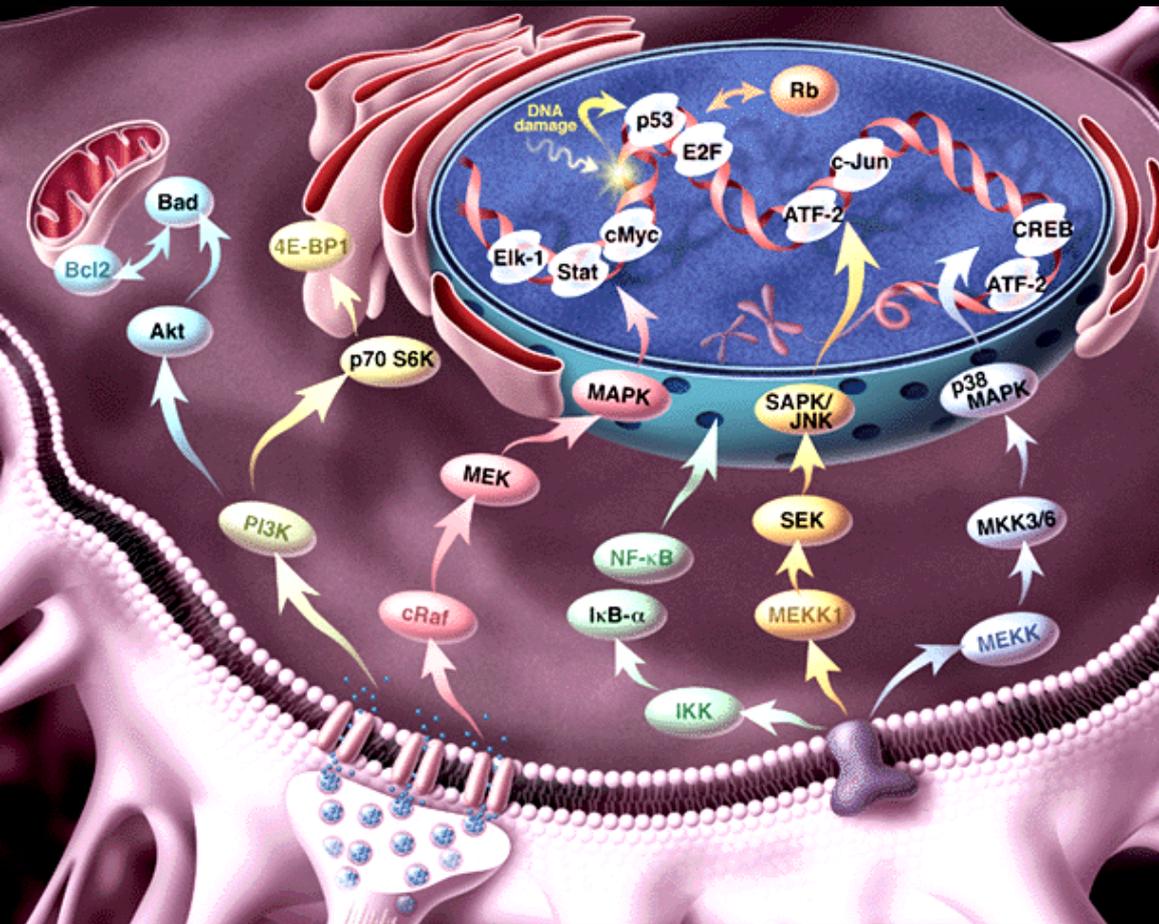
Multi-Color Flow Cytometry

- Combines complex immunophenotypic analysis with functional analysis (intracellular biochemical events)
 - Detailed characterization of rare subsets (e.g., antigen-specific T cells)
 - Identify new subsets more specifically associated with mechanism and clinical parameters
- Largely unappreciated in much of molecular biology– 11 parallel assays (classic) is NOT the same as 11 simultaneous measurements.

Single Cell Analysis in Drug Discovery

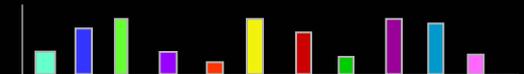


From phospho-molecular profiling to Signaling pathways



Raf
PKA
Erk
p38
PKC
PIP2
Jnk
PIP3
PIcy
Akt

Cell1



Cell2



Cell3



Cell4



Cell x



Signaling Pathway data

Single Cell Standards Issues to consider

- # of simultaneous parameters to measure
- Absolute vs relative (qualitative vs quantitative)
- Baseline fluorescence standards (quantitative)

- How does one FIND an informational 'blob' in n-space (feature extraction)?
- How does one represent such a 20-dimensional object?

- How does one apply such knowledge from flow cytometry to 3D imaging (confocal, other cytometry)?
- How does one deal with solid tissue slice (tissue biopsies)

- SOPs for sample handling?

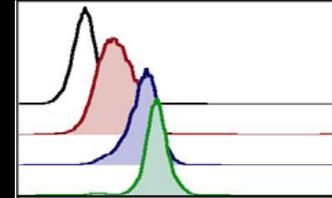
Why use Single Cells to Measure Cell events?

- Traditional



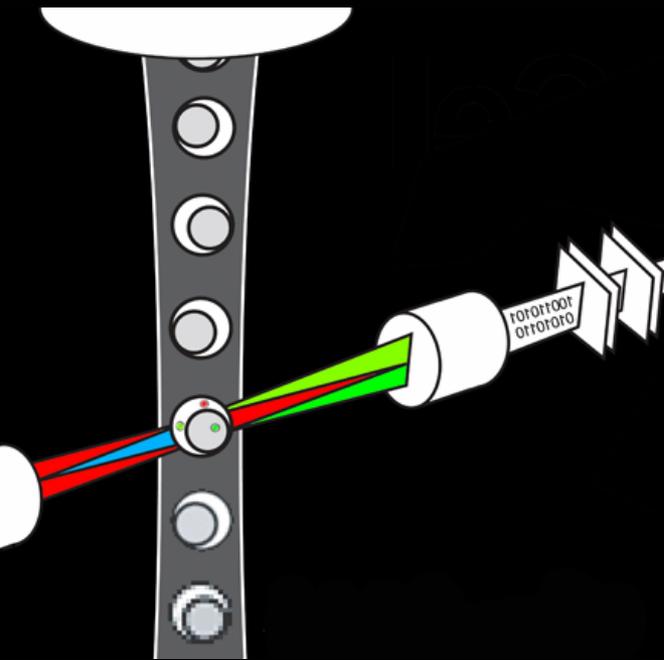
- Population analysis
- Homogeneous cell type: sorting, depletion

- Flow Cytometry



- Single cell analysis
- Heterogeneous populations can be separated via surface

Pulses to Numbers



- Advantages of new digital processing over Analog:
 - Highly accurate (acquisition time enhanced)
 - Can correct for dye spillover (matrix algebra for n colors)
 - Obtain pulse geometry metrics and time
 - Perform statistics on raw linear data

Stuff that impacts sensitivity

Fluorescence Sensitivity

Instrument

Q

- Optical design efficiency
- Laser power intensity
- PMT quantum efficiency
- Sheath flow rate

B

- Component autofluorescence
- Component scatter
- Raman scatter

Sample

Cell autofluorescence

Laser power intensity

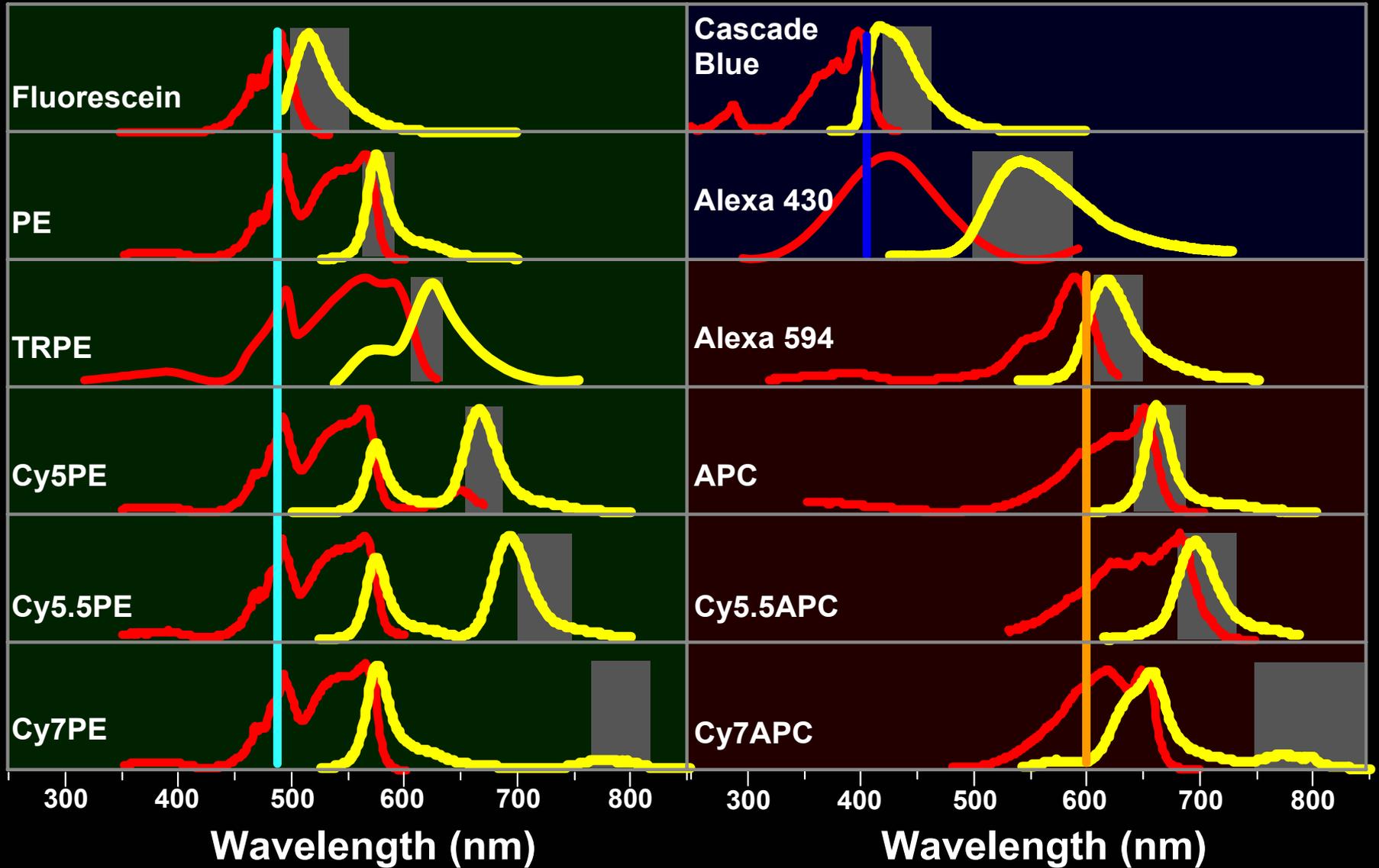
Unbound dye/fluorochrome

Laser power intensity
Sample flow rate

Spectral overlap

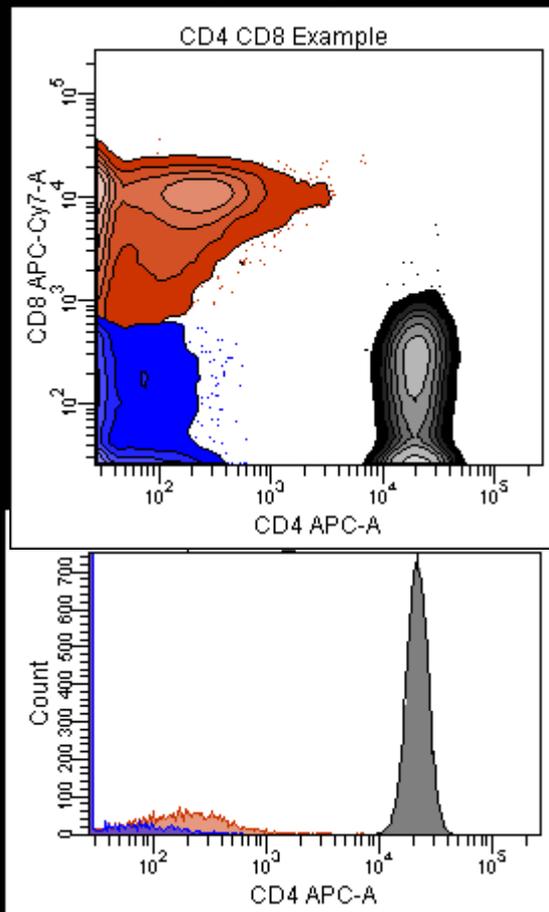
Laser power intensity
Filter design

Multi-color FACS: Spectral Overlap

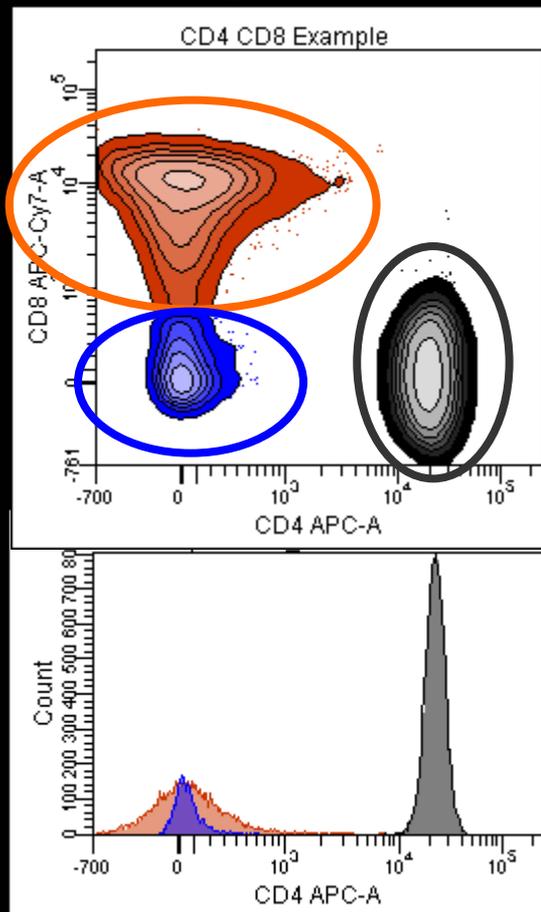


Stanford Biexponential Display (Logicle)

Log Display

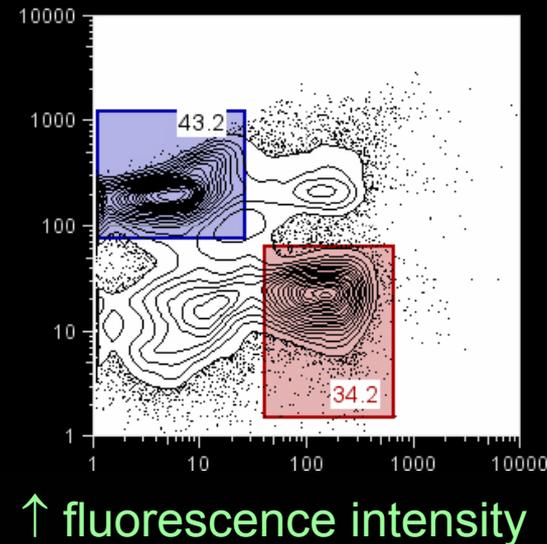
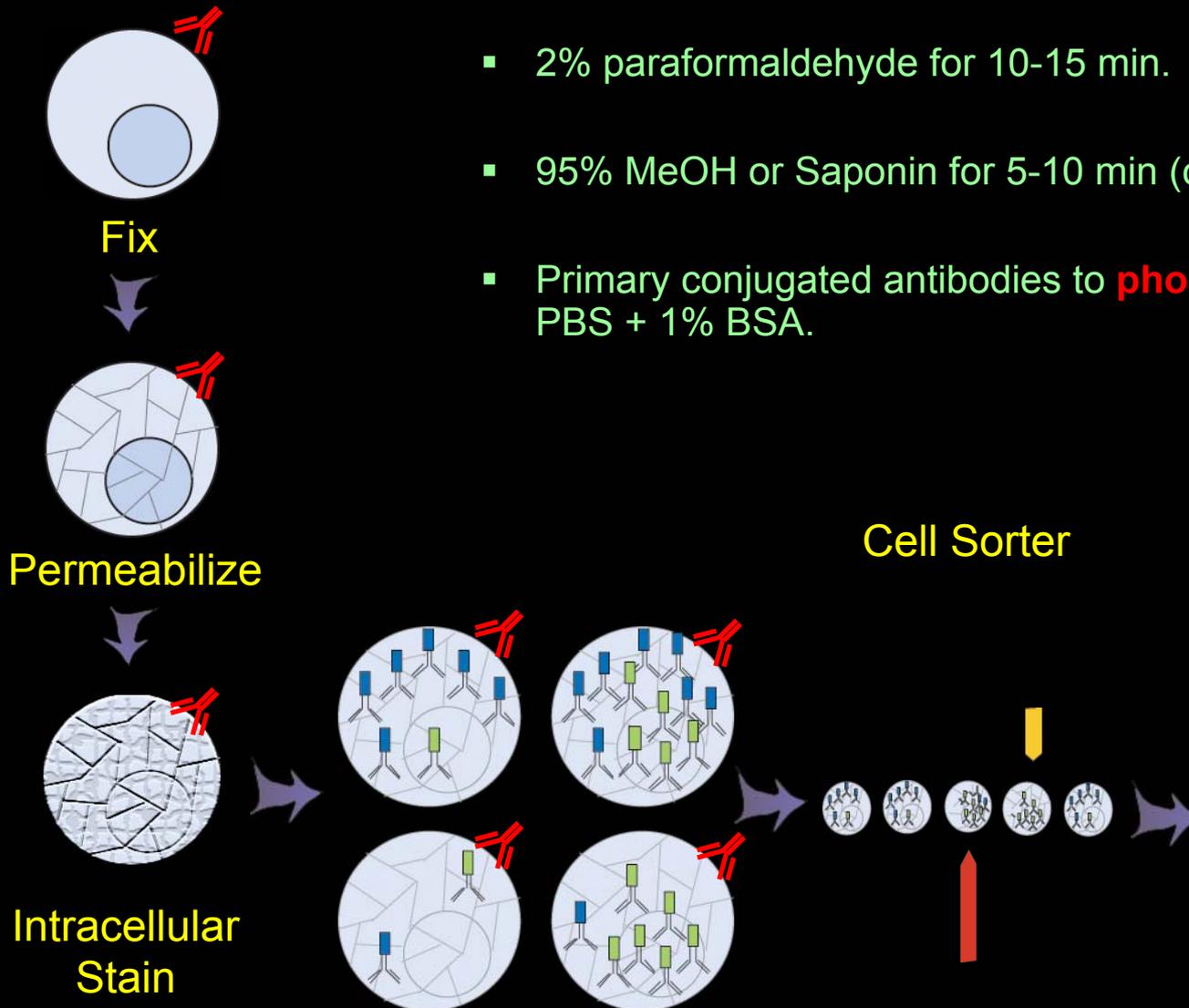


Biexponential



Intracellular Flow Cytometry Technique

- Stain with antibodies to surface proteins
- 2% paraformaldehyde for 10-15 min.
- 95% MeOH or Saponin for 5-10 min (cell type dependent).
- Primary conjugated antibodies to **phospho-epitopes** in PBS + 1% BSA.



Increasing Phospho Ab Repertoire

- Phospho Antibodies
- p38 MAPK
- JNK, cJun
- AKT, PIP2, PIP3,
- PKC $\alpha/\beta/\theta/\delta$, Rsk
- Raf, Mek, ERK, ELK
- Rsk, Creb,
- STAT1,3,5,6, c-Src
- CREB, cJUN, IKK α
- p53 s15, s20 s37, s392
- Pyk2, Shc, Fak, src
- Slp76, Zap70, Syk, Lat, Vav,
- Lck, PLC γ
- Beta-integrins

Every new antibody increases the potential of discovering entirely new correlations for disease processes (targets and diagnostics) as well as utility in drug design and development

Phospho Antibodies

- EGFR Pkg
- PDGFR RB
- cKit NFAT
- VEGFR NFKB
- PKA Caveolin
Paxillin

Stimulation of Murine Splenocytes

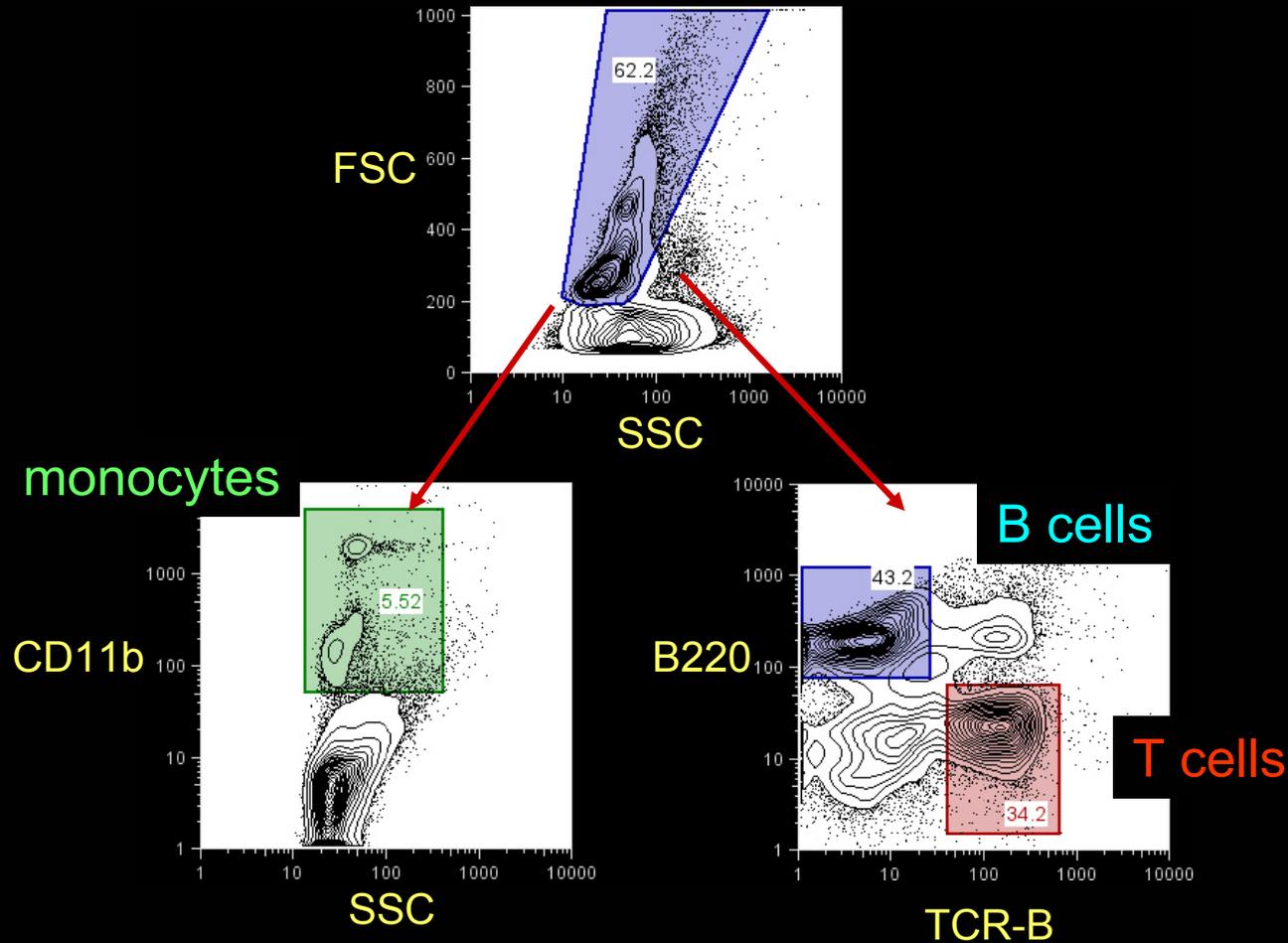
Dendritic Cell Subpopulation Analysis (B220⁻ CD8⁻ CD11c⁺)



Collect Splenic cells
10 Minutes post-
injection of IFN γ (in
vivo)

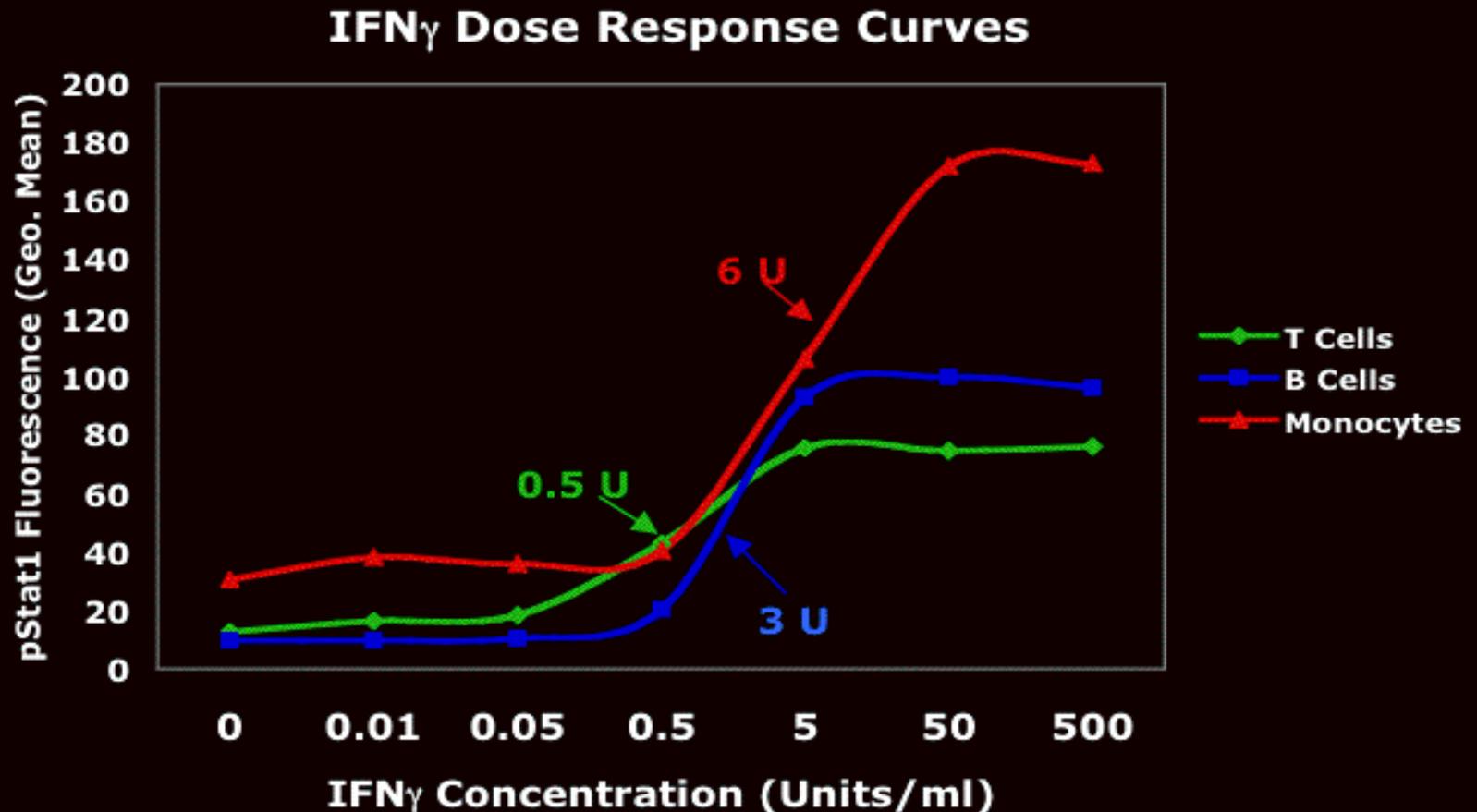
Read out Stat1
transcription factor
activation via its
phosphorylation

Murine Splenocytes - Gating



Phospho-FACS allows for Pharmacodynamics in Vivo

Cell Subset Specific IFN γ Sensitivity across a titration

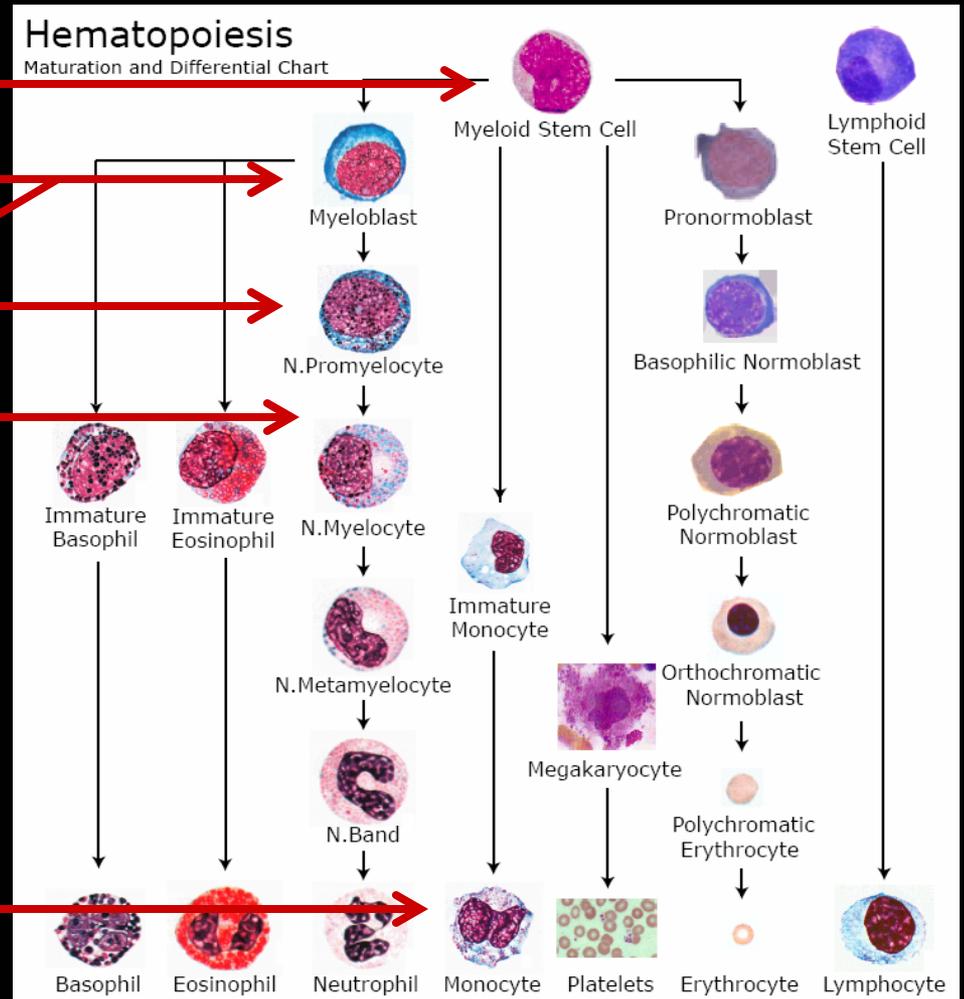


Leukemia (AML) Classification by Differentiation

- M0 – undifferentiated AML
- M1 – myeloblastic, immature
- M2 – myeloblastic, mature
- M3 – promyelocytic
- M4 – myelomonocytic
- M5 – monocytic

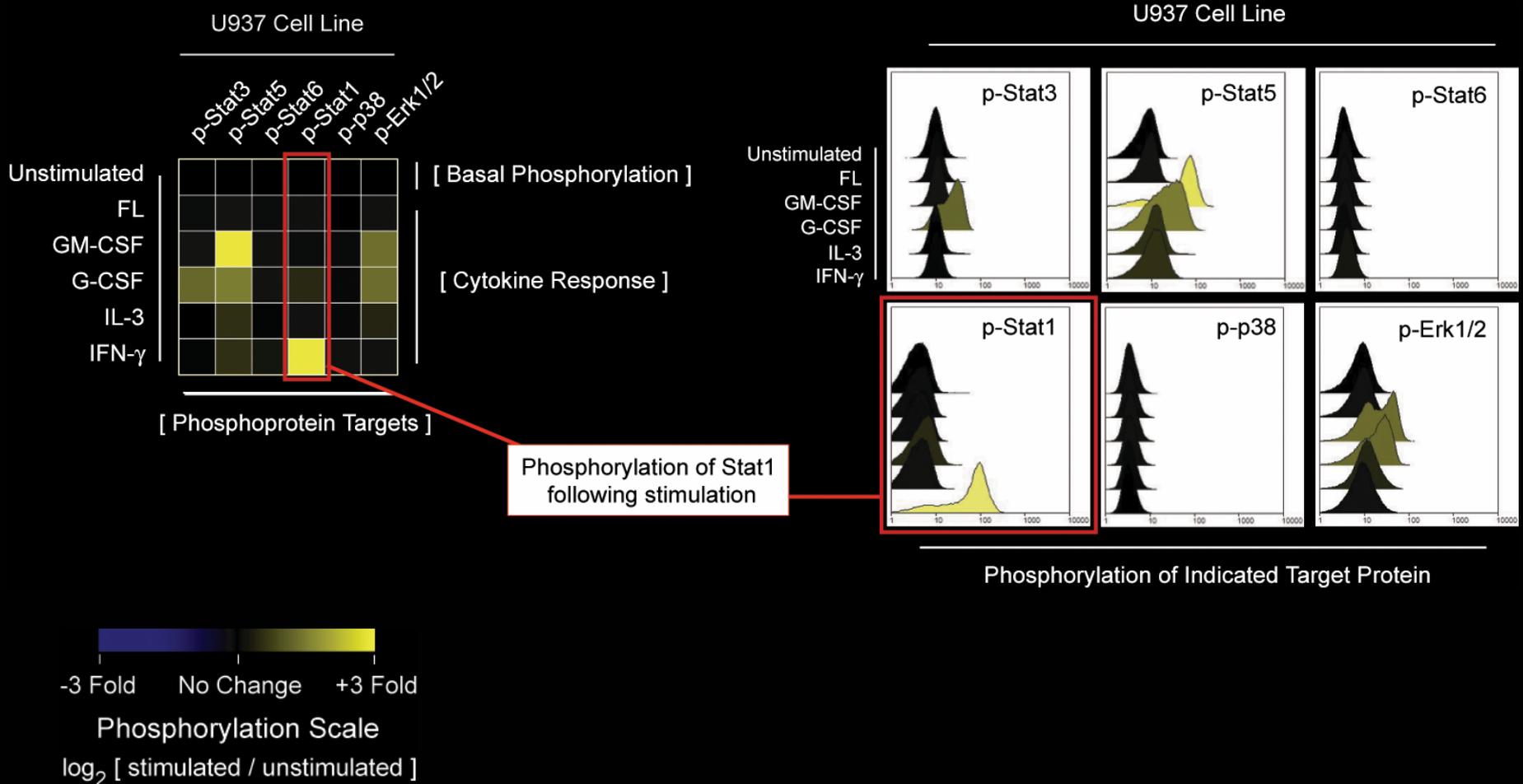
CD34 marker can be found on AMLs from all FAB classes

(lymphohematopoietic stem/progenitor cell marker)

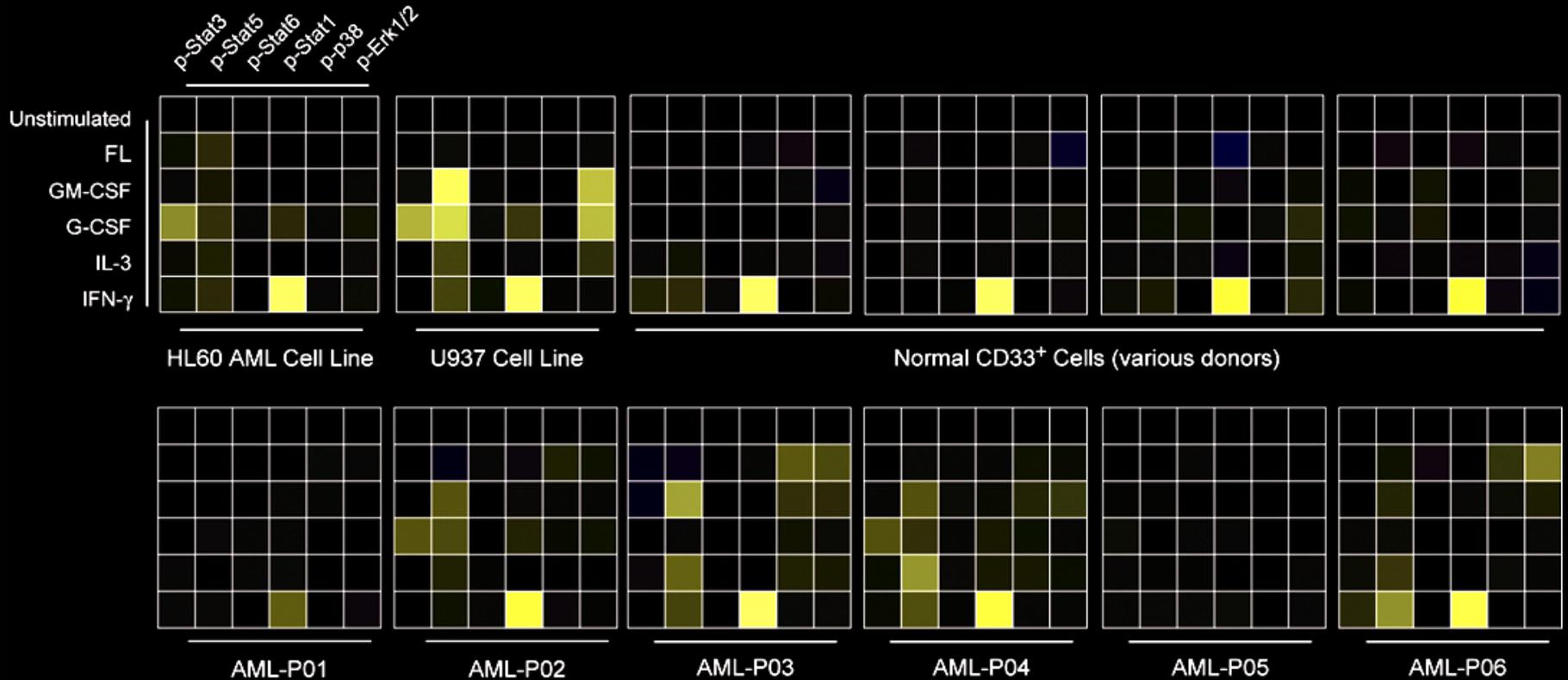


Could provoking cells to respond to external stimuli, such as cytokines, differentiate AML blasts with altered signal transduction networks?

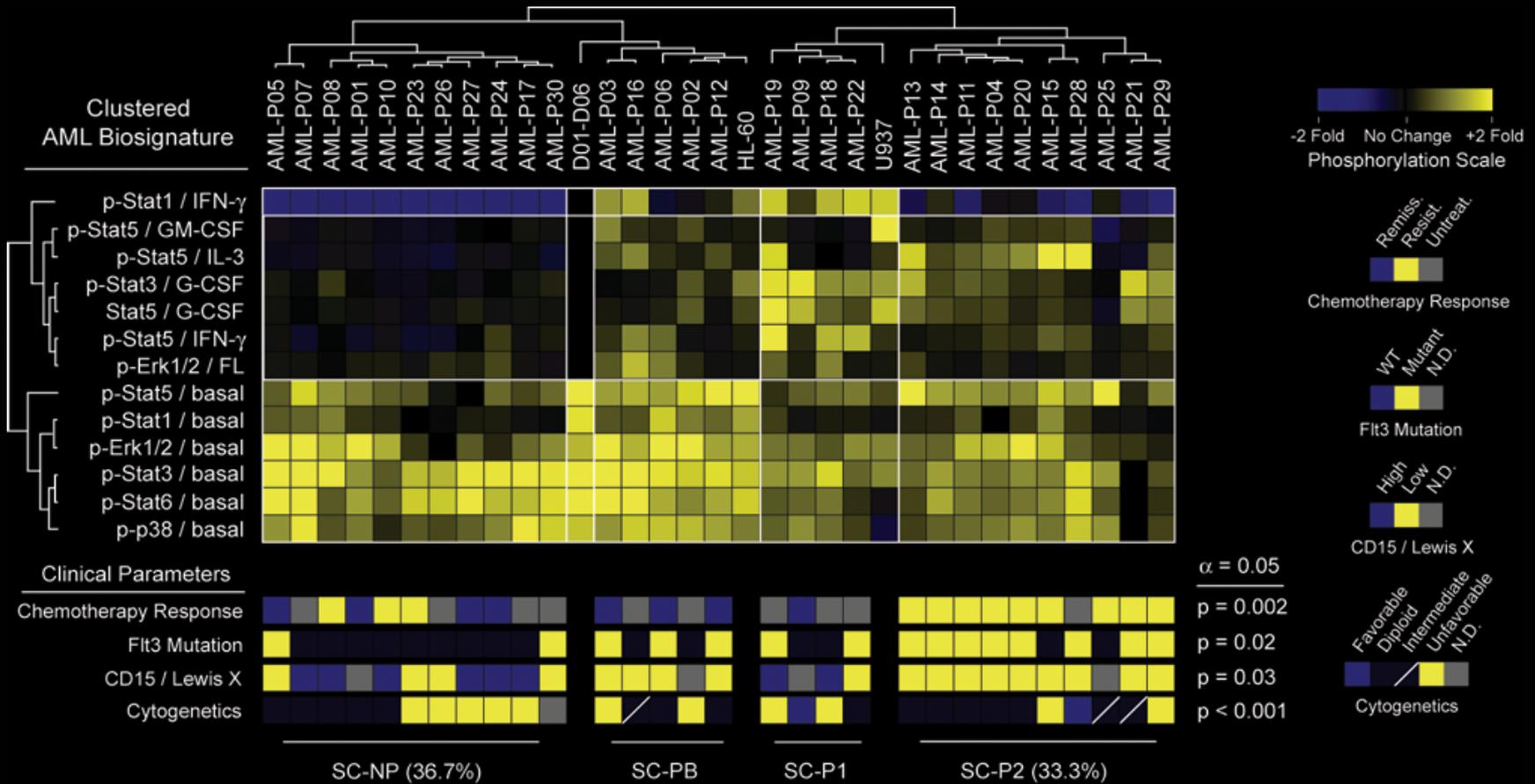
Model: Cytokine Response of U937 Cells



Cytokine Responses of Normal and Tumor Cells

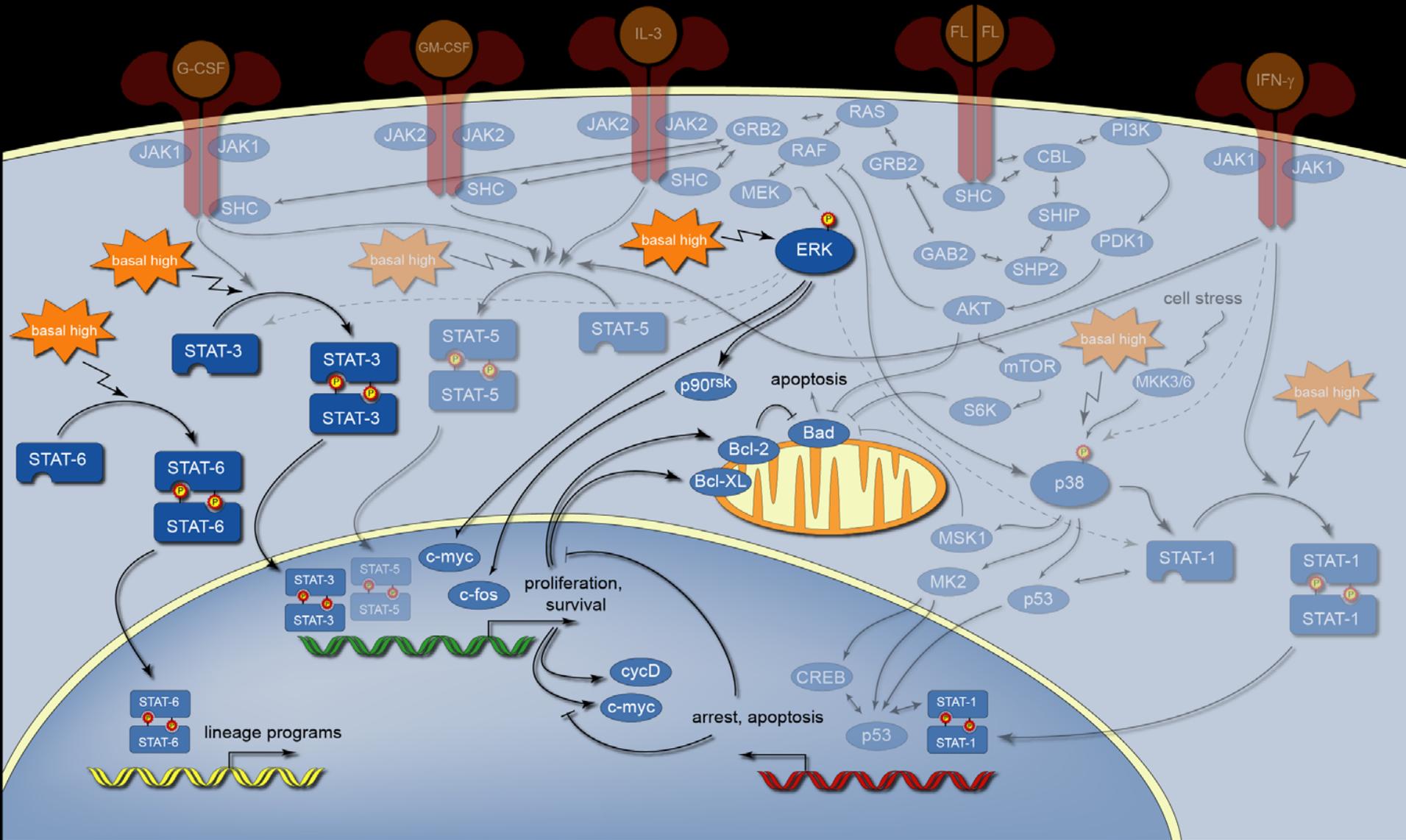


Clustering of Biosignature, Clinical Significance



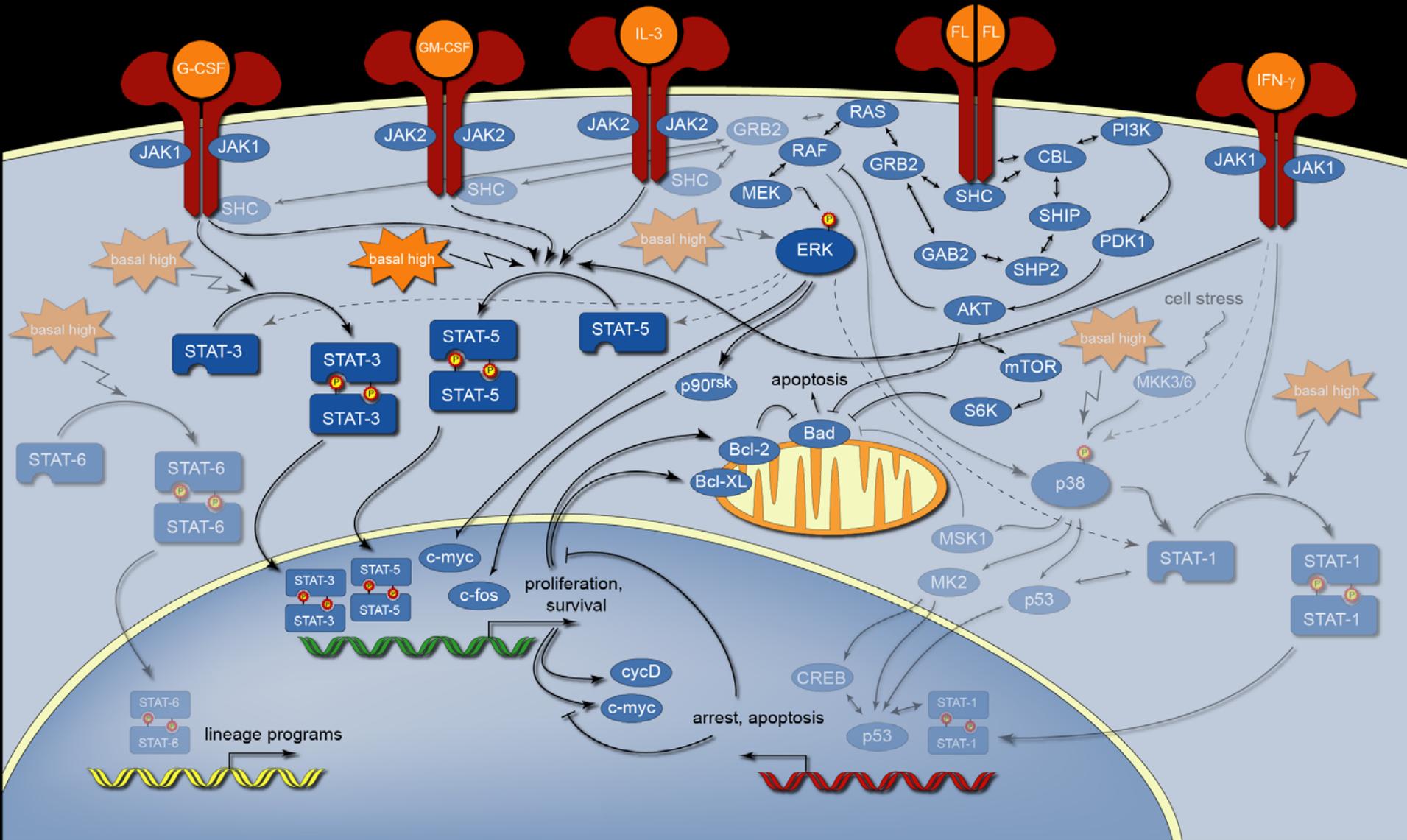
SC-NP (standard chemotherapy responses)

SC-NP Composite Profile



SC-P2 (Flt3 mutant, chemotherapy insensitive)

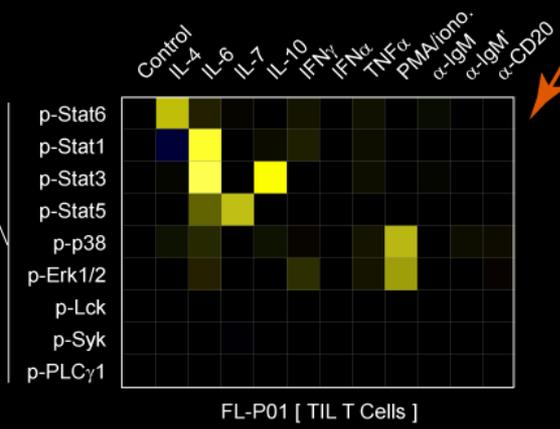
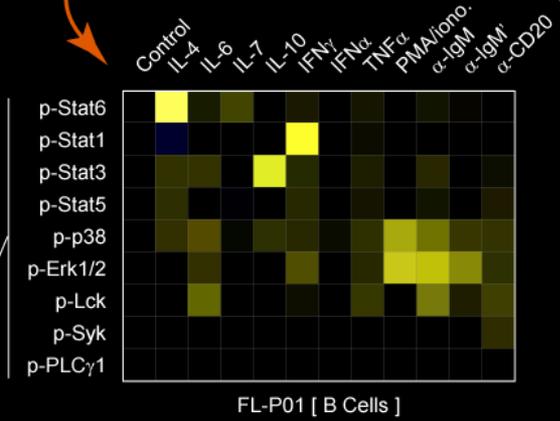
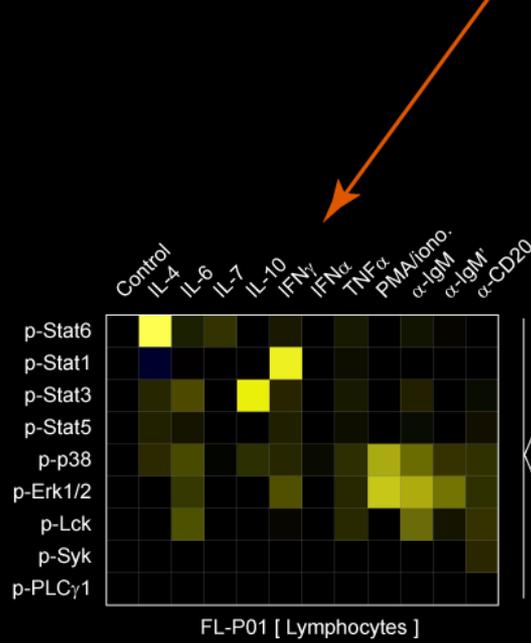
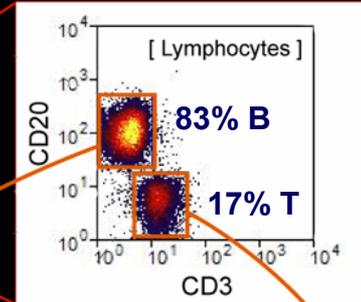
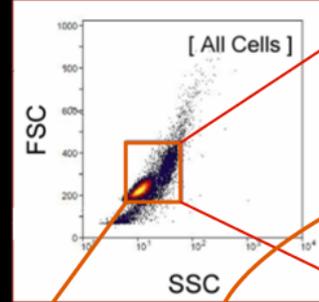
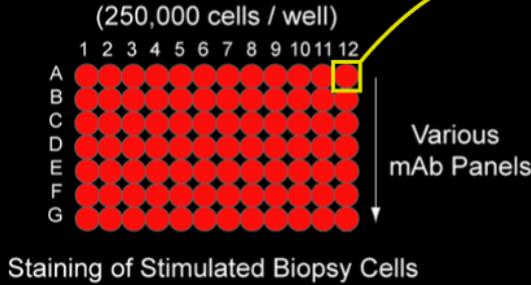
SC-P2 Composite Profile



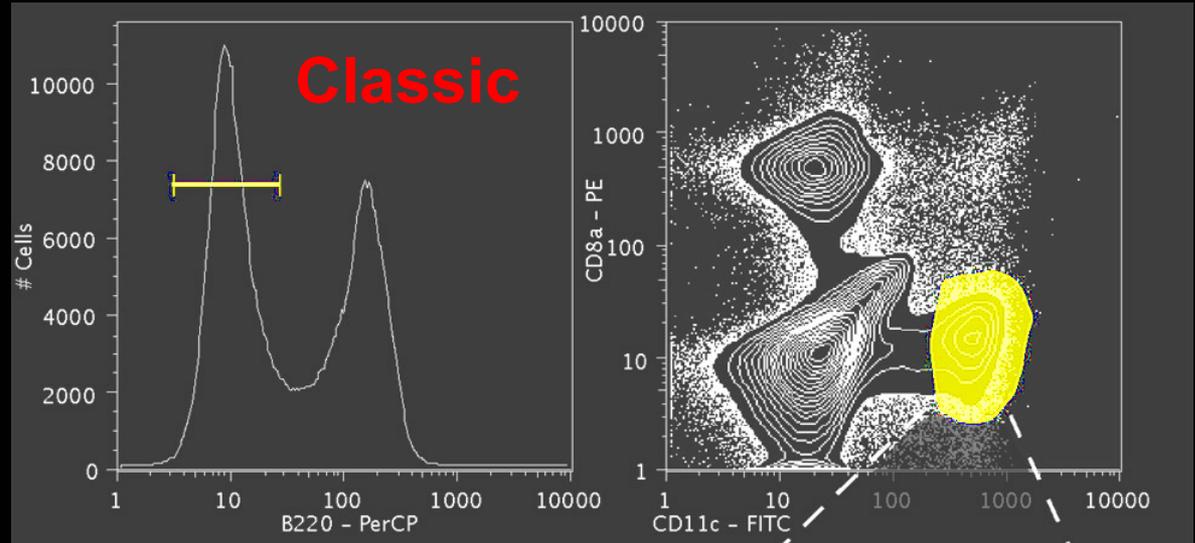
Array Overview of Lymphoma Signaling



Tumor Biopsy



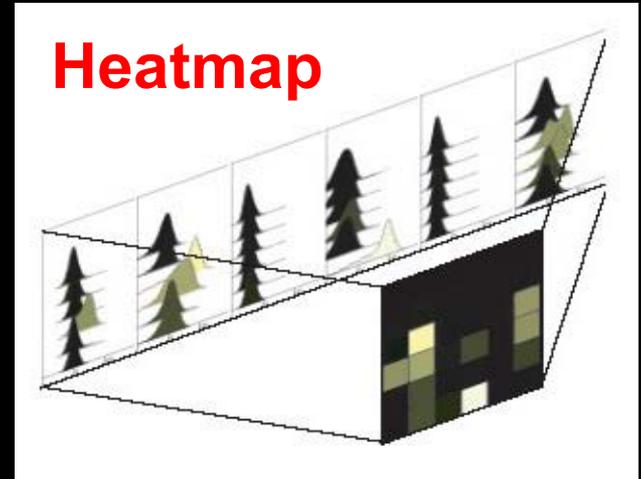
**New Approaches
To Representing
Single Cell Data
Present New Problems,
but suggest
Interesting possibilities**



Kinase 1
Kinase 2
Kinase 3
KinaseTarg 1
KinaseTarg 2
CD4
B220
Surf Marker X

**Multi-D
Single Cell**

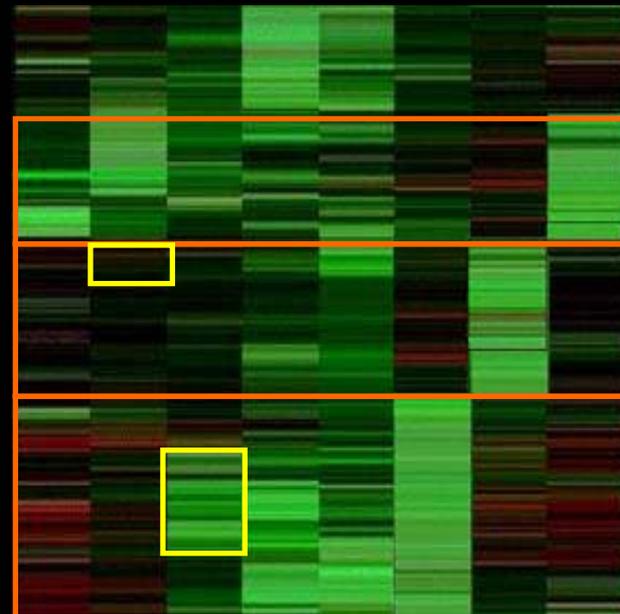
Heatmap



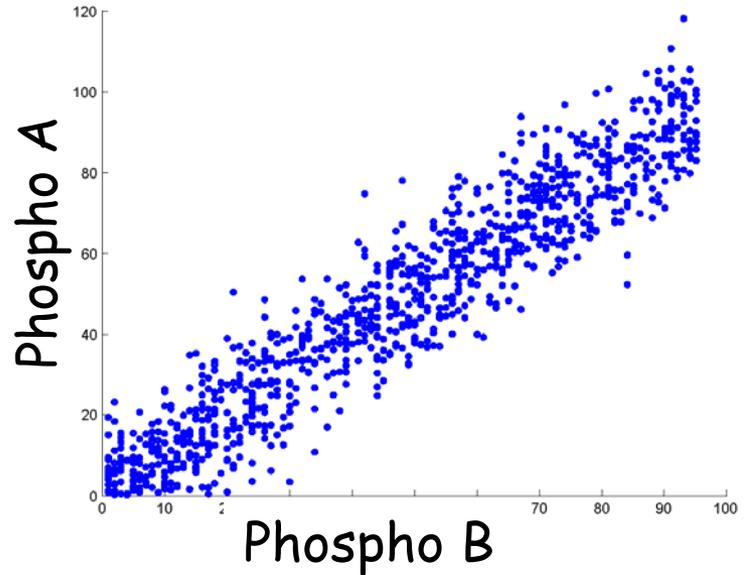
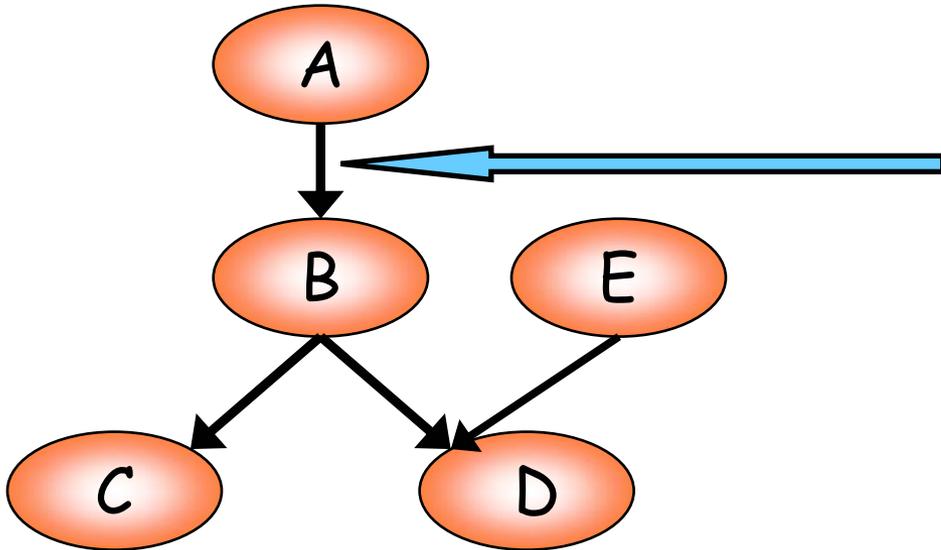
Macrophage

B cells

CD4+ T cells



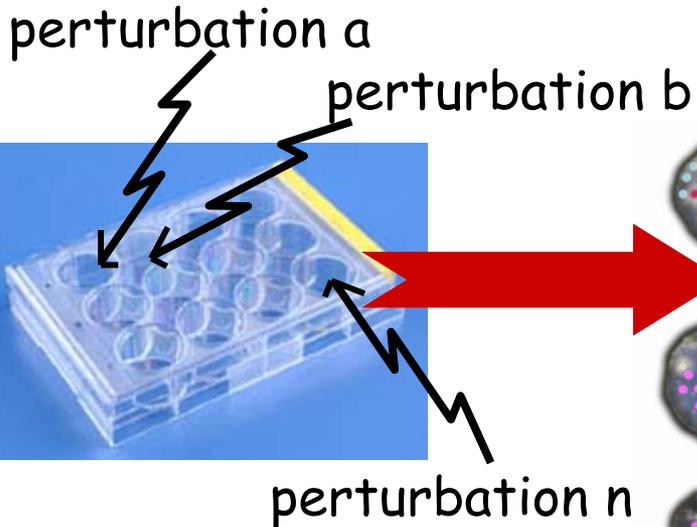
What is a Bayesian Network?



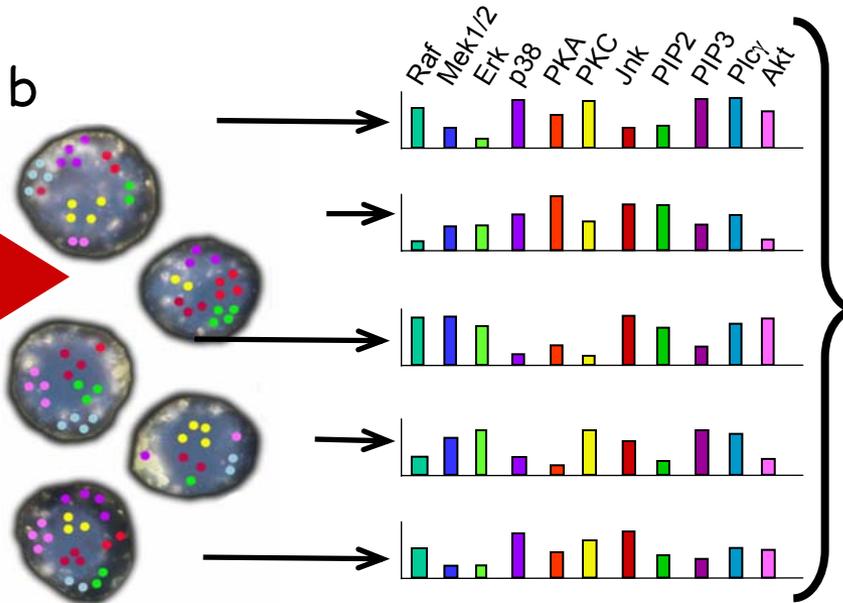
+ A Mathematical
(probabilistic) description
of the connections in the
graph ...

T-Lymphocyte Data

Conditions (96 well format)



11 Color Flow Cytometry



Datasets of cells

- *condition 'a'*
- *condition 'b'*
- *condition... 'n'*

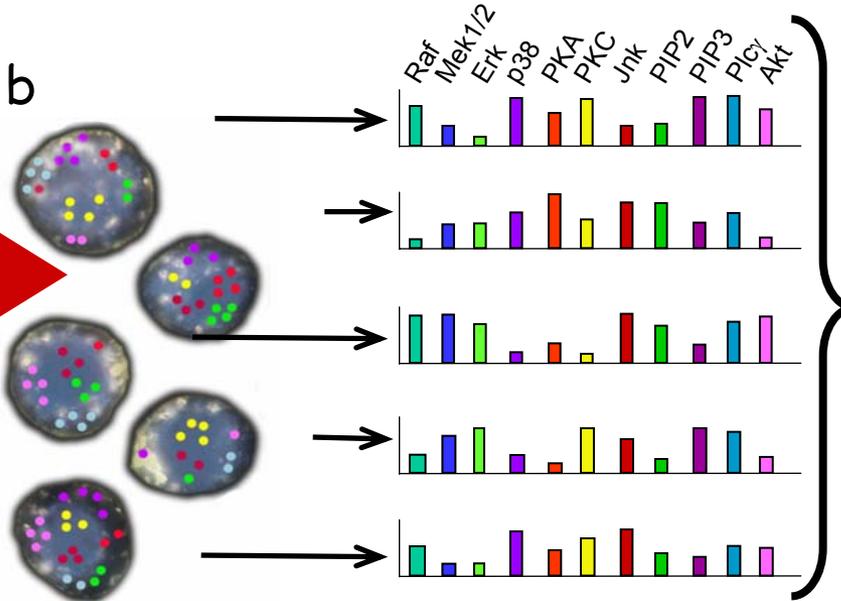
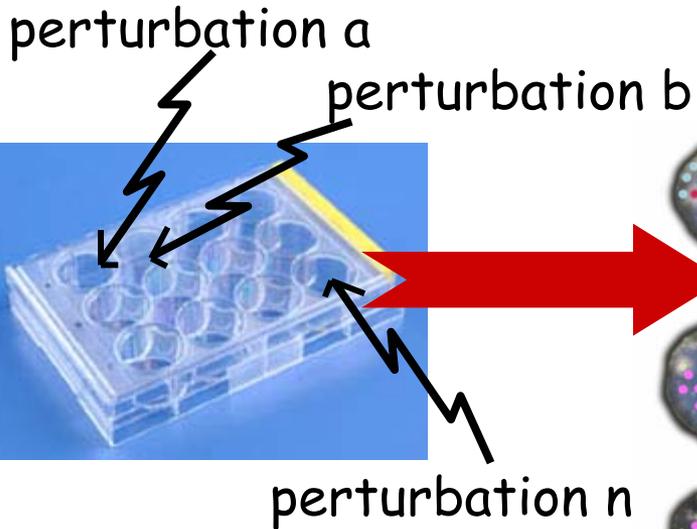
- Primary human T-Cells
- 9 conditions
 - (6 **Specific** interventions)

- 9 phosphoproteins, 2 phospholipids
- 600 cells per condition
 - 5400 data-points

T-Lymphocyte Data

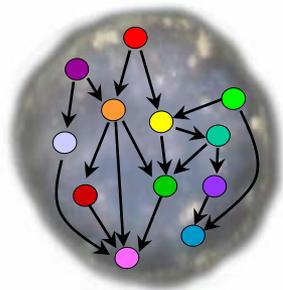
Conditions (96 well format)

11 Color Flow Cytometry



Datasets of cells

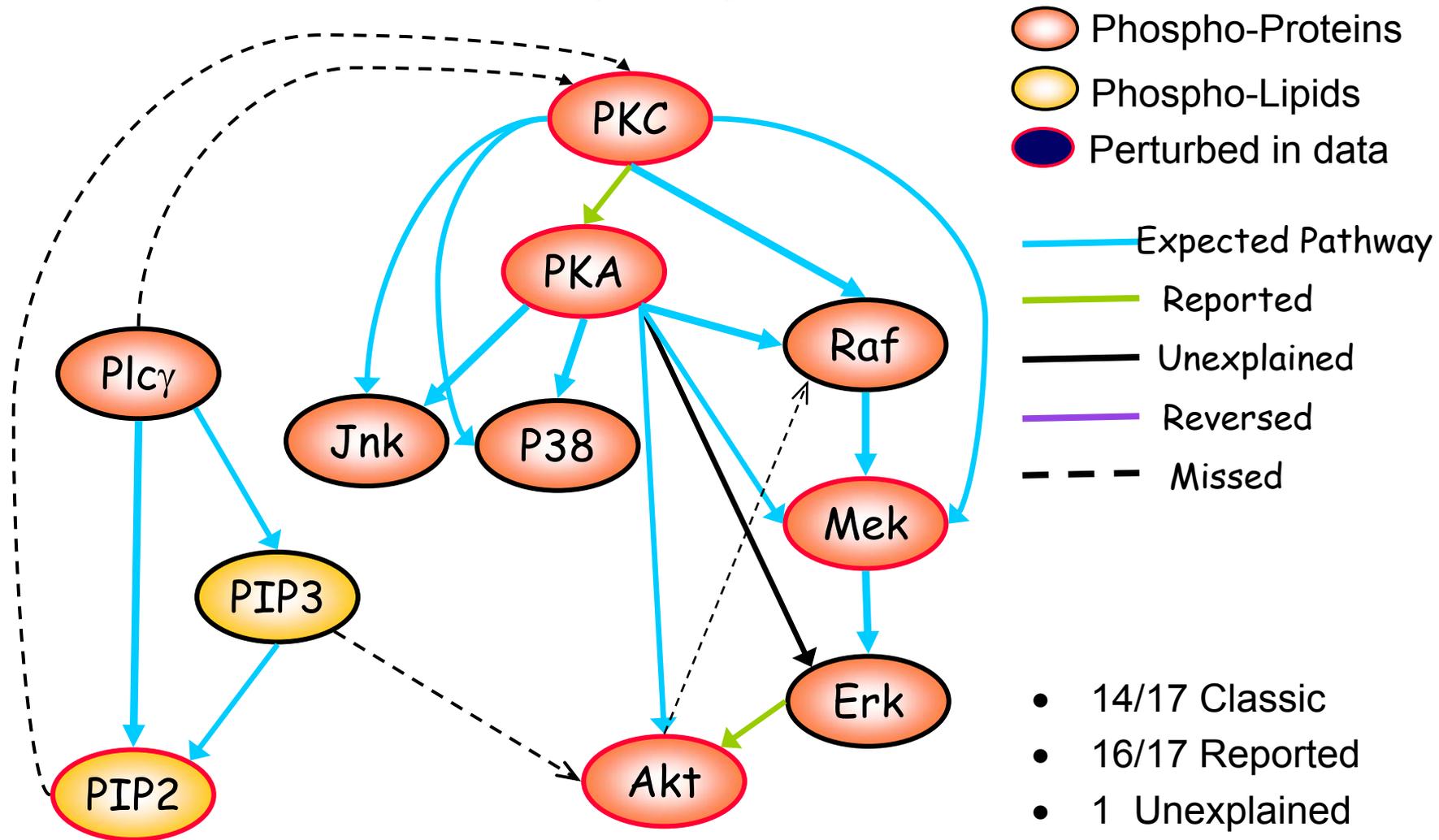
- condition 'a'
- condition 'b'
- condition... 'n'



Influence diagram of measured variables

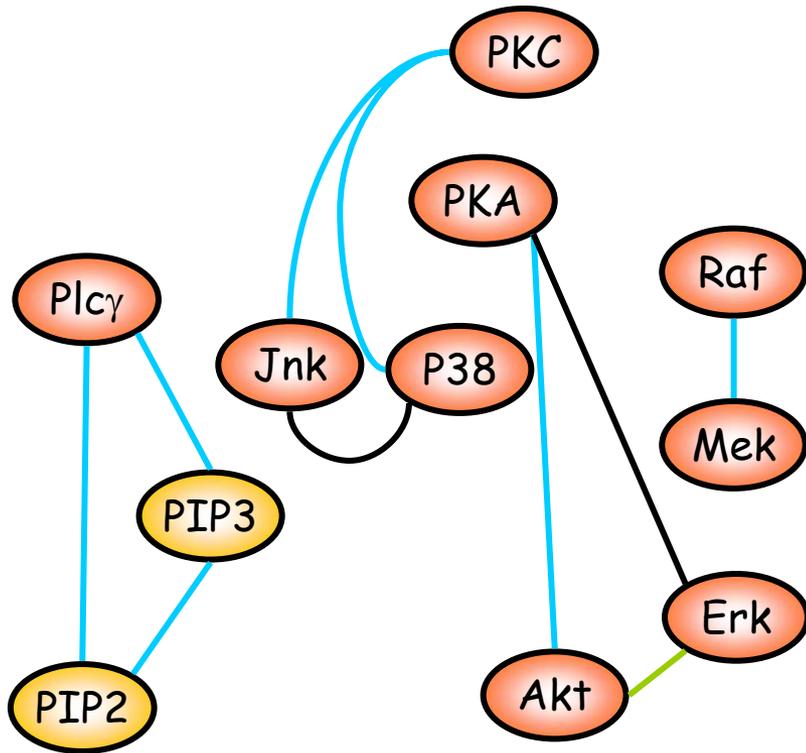
Bayesian Network Analysis

A T cell signaling map *ab initio* from multiparameter data by Bayesian Inference.



- 14/17 Classic
- 16/17 Reported
- 1 Unexplained
- 4 Missed

Interventions are Required for Directionality

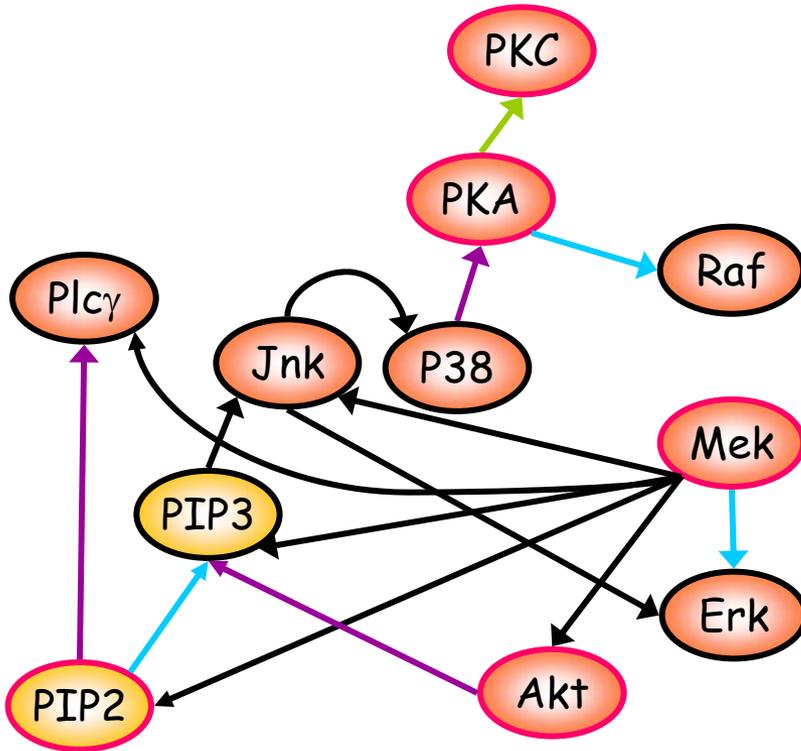


	Lacking Intervention data	Complete Dataset
<u>Expected</u>	7/10	14/17
<u>Reported</u>	1/10	2/17
<u>Reversed</u>	N/A	1
<u>Unexplained</u>	2	1
<u>Missed</u>	11	4

Dataset: 1200 samples:

- 2 conditions
- no interventions

Simulated Westerns Diminish Network Integrity



	"Western blot"	Complete Dataset
Expected <u> </u>	6/16	14/17
Reported <u> </u>	1/16	2/17
Reversed <u> </u>	3	1
Unexplained <u> </u>	8	1
Missed	12	4

Simulated western blot: 420 samples:

- 14 conditions
- Each point average of 20 random cells

Huge Problem with complex instrumentation

- Setting up the machine to ensure valid output.
- Setting up complex experiments in an automated fashion.
- ‘Forcing’ students/technical staff to conform.

*FacsXpert** and the *Libris DataStore*

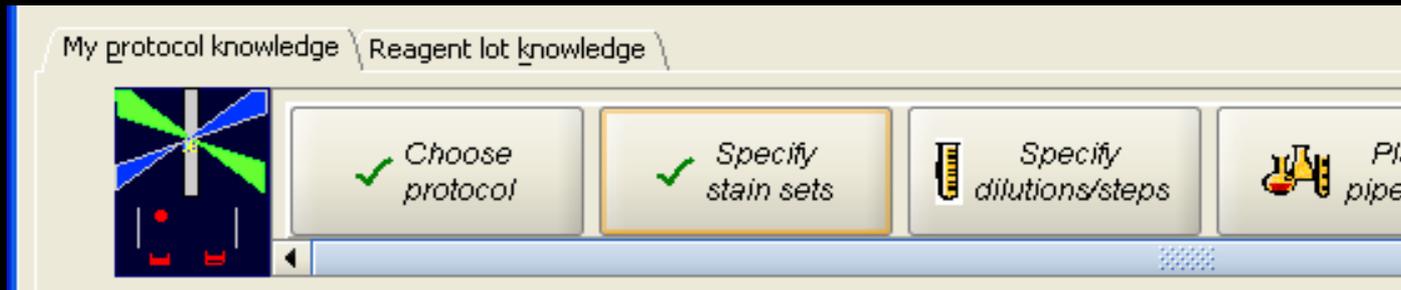
Designed to help researchers:

- Cope with this complexity when designing and executing FACS experiments
- Comply and with demanding requirements for long-term recoverability of FACS and other large data sets (Collaborative Electronic Notebook standards Association (CENSA)), US 21 CFAR part 11

**a knowledge-based system, Herzenberg laboratory (sold by ScienceXperts, Inc.)*

Start by choosing a new/existing protocol, specify

- Study and experiment name, Subject species, Cell source (tissue)



Take the individual through the experimental planning

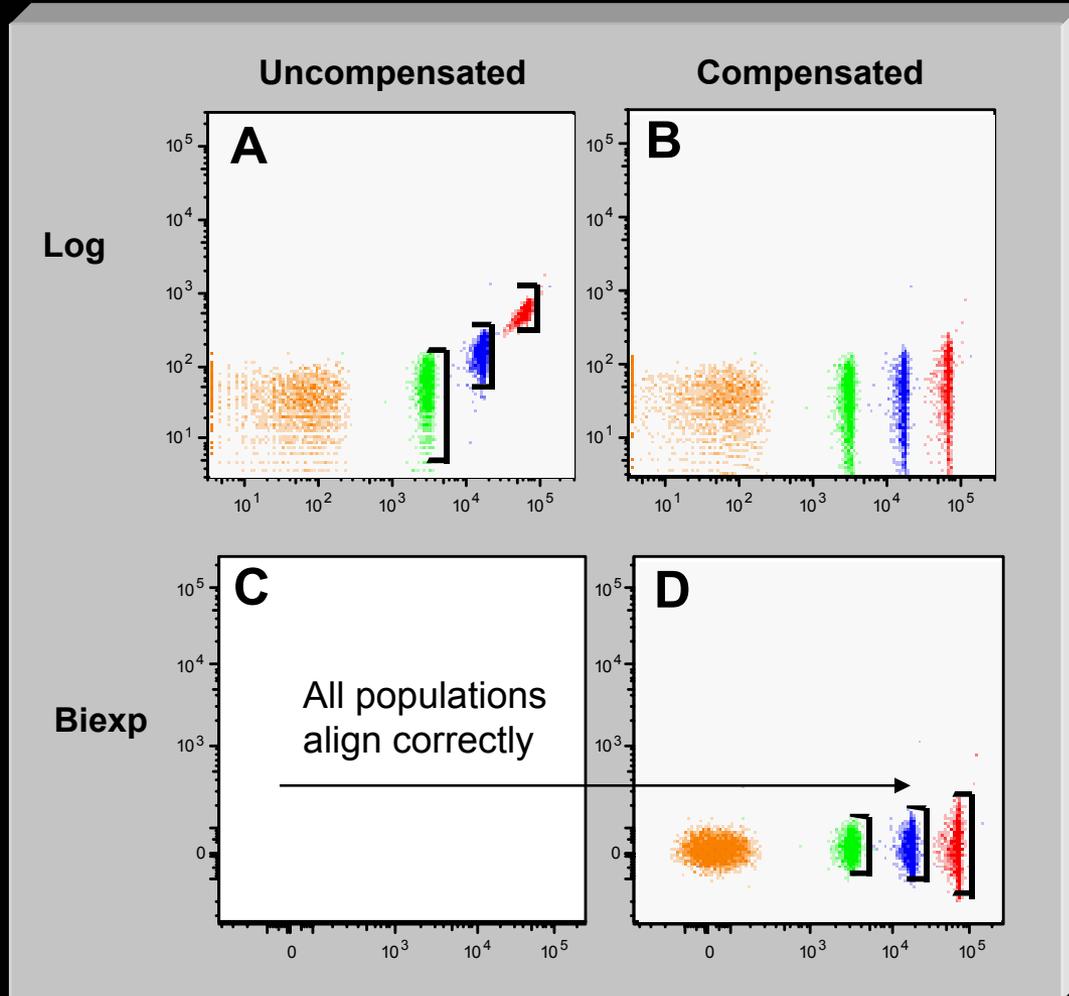


Carry out experiment, collect data, store, analyze



Important to validate instrument setup in an automated manner

APC-Cy7 Area



APC Area

Antibody capture beads stained with 3 levels of an APC reagent

The transformed display shows aligned populations in the APC-Cy7 dimension

Single Cells are an Unparalleled Information Resource... but...

- Common standards needed for instrument setup, runs.
- Automated experiment setup/protocols
 - *intelligent notebooks*
- Standards for representation of multi-D populations.
 - what is a population and what is the biological inference?
 - Cluster analysis
- Support (i.e. \$\$) for new visualization of multi-D

Acknowledgements



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Publications

- Perez O.D. et al. (2002). **Immunity**. 16:51-65.
- Perez O.D. and Nolan G.P. (2002) **Nature Biotechnology** 20:155-162.
- Krutzik PO, Irish J, Nolan GP, and Perez OD, Analysis of Phospho-Proteins by Flow Cytometry: Techniques and Clinical Applications. **Clinical Immunology Reviews**, 2003, December.
- Perez OD and Nolan GP. Flow cytometric analysis of kinase signaling cascades. **Methods in Flow Cytometry**. Humana Press. Editor: Howard Shapiro, 2004, March.
- Krutzik PO and Nolan GP. Intracellular phospho-protein staining techniques for flow cytometry: monitoring single cell signaling events. **Cytometry**, 2003, September
- Perez, OD and Nolan et al. LFA-1 lowers of T-cell activation thresholds and signaling through cytohesin-1 and JAB-1. **Nature Immunology**, 2003, November.
- Irish, J. and Nolan et al. Single cell profiling of potentiated phospho-protein networks in cancer cells. **Cell**, August, 2004.

<http://proteomics.stanford.edu>

<http://www.stanford.edu/group/nolan>